

Alpha Virus nsp4 alignment

CLUSTAL W (1.82) multiple sequence alignment

SFV_nsp4 PLTAEDKQDEDRRRAL SDEVSKWFR TGLGAELEV ALTSRYEV EGCKSILIAMTTLARDIK 594
RRV_nsp4 PLPAGDTQDEDRRRAL KDETDRWARVGLKSELEIA LSSRYEVNGTGNIVRAMATLAKSLK 594
ONV_nsp4 PLAAGDEQDEDRRRALADEVTRWQRTGLVTELEKAVY SRYEVQGITAVITSMATFANSKE 594
SinV_nsp4 PLPADDEQDEDRRRALLDET KAWFRVGITGTLAVAVTTRYEV DNITPVLLALRTFAQSKR 599
VEEV_nsp4 PLAVDDEHDDRRRALHEESTRWNRVGILPELCKAVESRYETVGT SIIIVMAMTTLASSVK 593
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SFV_nsp4 AFKKLRGPVIHLYGGPRLVR 614
RRV_nsp4 NFKKLRGPPIVHLYGGPK--- 611
ONV_nsp4 NFKKLRGPVVTLYGGPK--- 611
SinV_nsp4 AFQAIRGEIKHLYGGPK--- 616
VEEV_nsp4 SFSYLRGAPITLY----- 606
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